







<u>Supplementary figure S1:</u> Standard curve analyses for modified E and sgRNA assays. Ct = crossing point, and is given on the Y axis. Absolute copy number is given on the X axis. Triplicate data points are plotted for each dilution. Standard curves are composed of 1:10 serial dilutions of a synthetic standard consisting of either E gene or sgRNA sequences.

Supplementary file S1

The sequences of the synthetic oligonucleotides used for quantification are given here. Primer sequences are underlined, probe sequences given in bold.

E gene synthetic oligonucleotide standard:

 $5'g cattg tag ag\underline{acagg tacgt taatag ttaatag cgt} actag tt \pmb{acactag ccatccttactg cgcttcg} at \underline{tgtgtgcgtactgctgcaatatg} \\ ttaac-3'$

sgRNA synthetic oligonucleotide standard

 $5'g cattgt \underline{cgatctcttgtagatctgttctc} taatattgctagtt \underline{accatccttactgcgcttcg} at \underline{tgtgtgcgtactgctgcaatat} g ttaacc-3'$